

## ABSTRACT

A method of identifying a ligand specific for a structural isoform of a protein by the binding of the structural isoform to a ligand on a support, a direct positional transfer of the structural isoform and a control isoform between two or more different solid or semi-solid supports and detection of at least one of the isoforms on each support, thus allowing for subtractive identification techniques to be used to identify the subset of ligands, or a ligand, specific for the structural isoform.

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